AMENDMENTS TO THE CLAIMS

In the Claims:

- 1.-9. (Canceled)
- 10. (Previously presented) A method of screening a test compound for its ability to induce cytochrome P-450 3A4 (CYP3A4 gene expression comprising
 - i) contacting said test compound with a protein comprised of a ligand binding domain of human pregnane X receptor (hPXR) having the amino acid sequence 141-434 of SEQ ID NO:14, wherein the protein shares at least 96% amino acid sequence identity with the ligand binding domain of SEQ ID NO:14 and retains the sequence's ligand-binding function,
 - ii) determining whether said test compound selectively binds to the ligand binding domain of said protein; and
 - iii) determining whether a test compound that selectively binds to the ligand binding domain of said protein induces receptor binding to a response element in the CYP3A4 gene promoter and expression of a cytochrome P-450 3A4 monooxygenase enzyme.
 - 11.-24. (Canceled)
- 25. (Previously presented) The method according to claim 10, wherein the method is an in vitro assay.
 - 26. (Canceled)
- 27. (Previously presented) The method according to claim 10 wherein the protein shares at least 97% amino acid sequence identity with the ligand binding domain of SEQ ID NO: 14 and retains the sequence's ligand-binding function.

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- 28. (Previously presented) The method according to claim 10 wherein said protein has an amino acid sequence including amino acids 130 to 434 of SEQ ID NO: 14.
- 29. (Previously presented) The method according to claim 10 wherein said protein shares at least 98% amino acid sequence identity with the ligand binding domain of SEQ ID NO: 14 and retains the sequence's ligand-binding function.
- 30. (Previously presented) The method according to claim 10 wherein said protein bears a detectable label.

31.-33. (Canceled)

34. (Previously presented) The method according to claim 10 wherein the ligand-binding domain of an hPXR polypeptide is fused to a DNA binding domain of a non-hPXR polypeptide.

35.-36. (Canceled)

- 37. (Previously presented) The method according to claim 25 wherein binding is determined by separating test compound bound to protein from free test compound and free protein.
- 38. (Currently amended) The method according to claim 10 wherein binding is determined by a scintillation proximity ass[[0]]ay.
- 39. (Previously presented) The method according to claim 10 wherein binding is determined by competitive bind assay.
 - 40. (Canceled)

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- 41. (Previously presented) A method of screening a test compound for its ability to bind to a protein comprising human pregnane X receptor ligand binding domain, thereby indicating an increased likelihood that the test compound alters in vivo expression of a cytochrome P-450 3A4 (CYP3A4) monooxygenase enzyme comprising:
 - i) contacting said test compound with a protein comprised of a ligand binding domain of human pregnane X receptor (hPXR), said ligand binding domain including the amino acid sequence of amino acids 141-434 of SEQ ID NO:14, wherein the protein comprises a domain sharing an amino acid sequence at least 96% identical to the ligand binding domain of SEQ ID NO:14, and
 - ii) determining whether said test compound selectively binds to the ligand binding domain of said protein.
- 42. (Currently Amended) The method according to claim 39, wherein a test compound of formula 1 is detectably labeled

$$\begin{array}{c|c} \text{Me}_3\text{C} & \text{H} & \text{O} \\ \text{HO} & \text{OR}_2 \\ \text{OME}_3 & \text{OR}_4 \end{array}$$

$$\begin{array}{c|c} \text{Me}_3\text{C} & \text{H} & \text{O} \\ \text{P} & \text{OR}_1 \\ \text{HO} & \text{OR}_2 \\ \text{CME}_3 & \text{OR}_4 \end{array}$$

and each of R1, R2, R3 AND R4 is, independently, C1-C6 alkyl (linear or branched).

43. (Currently Amended) A method for identifying a compound as an hPXR agonist, the method comprising:

providing a polypeptide comprising the ligand-binding domain of an hPXR, wherein the ligand-binding domain comprises amino acids 130-434 of SEQ ID NO:14, wherein the polypeptide selectively binds a detectably labeled compound of formula 1

$$\begin{array}{c|c} \text{Me}_3\text{C} & \text{H} & \text{O} \\ \text{HO} & \text{OR}_2 \\ \text{OME}_3 & \text{OR}_4 \end{array}$$

$$\begin{array}{c|c} \text{Me}_3\text{C} & \text{H} & \text{O} \\ \text{P} & \text{OR}_1 \\ \text{HO} & \text{OR}_2 \\ \text{CME}_3 & \text{OR}_4 \\ \end{array}$$

and each of R1, R2, R3 AND R4 is, independently, C1-C6 alkyl (linear or branched);

contacting the polypeptide with a test compound;

determining whether the binding of the polypeptide to the detectably labeled compound of formula 1 is altered in the presence of the test compound, a decrease in the binding being an indication that the test compound is a competitive inhibitor of the detectably labeled compound of formula 1; and

determining whether expression of a CYP3A4 gene product, following receptor binding to a response element in the CYP3A4 gene promoter, is altered in a cell in the presence of the test compound, wherein an increase in the expression is an indication that the test compound is useful as an hPXR agonist in screening assays.

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- 44. (Previously presented) The method according to claim 42 or 43, wherein the detectably labeled compound of formula 1 is GW-485801.
- 45. (Previously presented) The method according to claim 43, wherein the cytochrome P450 3A4 gene product is a cytochrome P-450 3A4 monooxygenase enzyme.
- 46. (Previously presented) The method of claim 10, wherein the ligand-binding domain is a fragment of SEQ ID NO:14 at least 75 consecutive amino acid residues in length.
- 47. (Previously presented) The method of claim 10, wherein the ligand-binding domain is a fragment of SEQ ID NO:14 at least 50 consecutive amino acid residues in length.
- 48. (Previously presented) The method of claim 10, wherein the ligand-binding domain is a fragment of SEQ ID NO:14 at least 30 consecutive amino acid residues in length.